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Reviewer: Anne Corrigan

Timestamp: [year=2009; month=1; day=2; hr=10; min=55; sec=54; ms=487;]

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Application No: 10537228

Version No: 3.0

Input Set:

Output Set:

Started: 2008-12-19 15:45:51.487

Finished: 2008-12-19 15:45:54.907

Elapsed: 0 hr(s) 0 min(s) 3 sec(s) 420 ms

Total Warnings: 21

Total Errors: 0

No. of SeqIDs Defined: 74

Actual SeqID Count: 74

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Input Set:

Output Set:

Started: 2008-12-19 15:45:51.487
Finished: 2008-12-19 15:45:54.907
Elapsed: 0 hr(s) 0 min(s) 3 sec(s) 420 ms
Total Warnings: 21
Total Errors: 0
No. of SeqIDs Defined: 74
Actual SeqID Count: 74

Error code

Error Description

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Ala Ser Leu Ser Ile Pro Val Ser Arg Gly Leu Pro Gln Gln Ser Ser
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Pro Gln Gln Leu Leu Ser Leu Gln Gly Leu His Ser Thr Ser Leu Leu
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Asn Gly Pro Met Leu Gln Arg Ala Leu Leu Leu Gln Gln Leu Gln Gly
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Leu Asp Gln Phe Ala Met Pro Pro Ala Thr Tyr Asp Gly Ala Ser Leu
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Thr Met Pro Thr Ala Thr Leu Gly Asn Leu Arg Ala Phe Asn Val Thr
115 120 125

Ala Pro Ser Leu Ala Ala Pro Ser Leu Thr Pro Pro Gln Met Val Thr
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Gly Pro Pro Pro Val Gly Val Pro Ile Asn Pro Ser Gln Leu Asn His
165 170 175

Ser Gly Arg Asn Thr Gln Lys Gln Ala Arg Thr Pro Ser Ser Thr Thr
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Pro Asn Arg Lys Asp Ser Ser Ser Gln Thr Val Pro Leu Glu Asp Arg
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Glu Asp Pro Thr Glu Gly Ser Glu Glu Ala Thr Glu Leu Gln Met Asp
210 215 220

Thr Cys Glu Asp Gln Asp Ser Leu Val Gly Pro Asp Ser Met Leu Ser
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Glu Pro Gln Val Pro Glu Pro Glu Pro Phe Glu Thr Leu Glu Pro Pro
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Ala Lys Arg Cys Arg Ser Ser Glu Glu Ser Thr Glu Lys Gly Pro Thr
260 265 270

Gly Gln Pro Gln Ala Arg Val Gln Pro Gln Thr Gln Met Thr Ala Pro
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Lys Gln Thr Gln Thr Pro Asp Arg Leu Pro Glu Pro Pro Glu Val Gln

290

295

300

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Lys Leu Leu Arg Gln Ala Gln Thr Gln Thr Ser Pro Glu His Leu Ala
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Gln Leu Gln Pro Arg Glu Thr Asp Pro Pro Asn Gln Ala Gln Ala Gln
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Thr Gln Pro Gln Pro Leu Trp Gln Ala Gln Ser Gln Lys Gln Ala Gln
370 375 380

Thr Gln Ala His Pro Gln Val Pro Thr Gln Ala Gln Ser Gln Glu Gln
385 390 395 400

Thr Ser Glu Lys Thr Gln Asp Gln Pro Gln Thr Trp Pro Gln Gly Ser
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Val Pro Pro Pro Glu Gln Ala Ser Gly Pro Ala Cys Ala Thr Glu Pro
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Gln Leu Ser Ser His Ala Ala Glu Ala Gly Ser Asp Pro Asp Lys Ala
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Leu Pro Glu Pro Val Ser Ala Gln Ser Ser Glu Asp Arg Ser Arg Glu
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Ala Ser Ala Gly Gly Leu Asp Leu Gly Glu Cys Glu Lys Arg Ala Gly
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Leu Gln Ser Ser Asn Ser Arg Ala Phe Asn Thr Thr Pro Leu Thr Ser
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Gly Pro Arg Pro Gly Asp Ser Thr Ser Ala Thr Pro Ala Ile Ala Ser
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Thr Pro Ser Lys Gln Ser Leu Gln Phe Phe Cys Tyr Ile Cys Lys Ala
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Ser Ser Ser Ser Gln Gln Glu Phe Gln Asp His Met Ser Glu Ala Gln
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His Gln Gln Arg Leu Gly Glu Ile Gln His Ser Ser Gln Thr Cys Leu
565 570 575

Leu Ser Leu Leu Pro Met Pro Arg Asp Ile Leu Glu Lys Glu Ala Glu
580 585 590

Asp Pro Pro Pro Lys Arg Trp Cys Asn Thr Cys Gln Val Tyr Tyr Val
595 600 605

Gly Asp Leu Ile Gln His Arg Arg Thr Gln Glu His Lys Val Ala Lys
610 615 620

Gln Ser Leu Arg Pro Phe Cys Thr Ile Cys Asn Arg Tyr Phe Lys Thr
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Pro Arg Lys Phe Val Glu His Val Lys Ser Gln Gly His Lys Asp Lys
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Ala Gln Glu Leu Lys Thr Leu Glu Lys Glu Thr Gly Ser Pro Asp Glu
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Asp His Phe Ile Thr Val Asp Ala Val Gly Cys Phe Glu Ser Gly Gln
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Glu Ala Glu Glu Glu Phe Cys Lys Gln Val Lys Pro Arg Glu Thr Ser
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Ser Glu Gln Gly Lys Gly Ser Glu Thr Tyr Asn Pro Asn Thr Ala Tyr
725 730 735

Gly Glu Asp Phe Leu Val Pro Val Met Gly Tyr Val Cys Gln Ile Cys
740 745 750

His Lys Phe Tyr Asp Ser Asn Ser Glu Leu Arg Leu Ser His Cys Lys
755 760 765

Ser Leu Ala His Phe Glu Asn Leu Gln Lys Tyr Lys Ala Lys Asn Pro
770 775 780

Ser Pro Pro Pro Thr Arg Pro Val Ser Arg Lys Cys Ala Ile Asn Ala
785 790 795 800

Arg Asn Ala Leu Thr Ala Leu Phe Thr Ser Ser His Gln Pro Ser Pro
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Ala Phe Asn Val Thr Ala Pro Ser Leu Ala Ala Pro Ser Leu Thr Pro
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Pro Gln Met Val Thr Pro Asn Leu Gln Gln Phe Phe Pro Gln Ala Thr
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Arg Gln Ser Leu Leu Gly Pro Pro Pro Val Gly Val Pro Ile Asn Pro
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Ser Gln Leu Asn His Ser Gly Arg Asn Thr Gln Lys Gln Ala Arg Thr

100

105

110

Pro Ser Ser Thr Thr Pro Asn Arg Lys Thr Val Pro Leu Glu Asp Arg
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Glu Asp Pro Thr Glu Gly Ser Glu Glu Ala Thr Glu Leu Gln Met Asp
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Thr Cys Glu Asp Gln Asp Ser Leu Val Gly Pro Asp Ser Met Leu Ser
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Glu Pro Gln Val Pro Glu Pro Glu Pro Phe Glu Thr Leu Glu Pro Pro
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Ala Lys Arg Cys Arg Ser Ser Glu Glu Ser Thr Glu Lys Gly Pro Thr
 180 185 190

Gly Gln Pro Gln Ala Arg Val Gln Pro Gln Thr Gln Met Thr Ala Pro
 195 200 205

Lys Gln Thr Gln Thr Pro Asp Arg Leu Pro Glu Pro Pro Glu Val Gln
 210 215 220

Met Leu Pro Arg Ile Gln Pro Gln Ala Leu Gln Ile Gln Thr Gln Pro
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Lys Leu Leu Arg Gln Ala Gln Thr Gln Thr Ser Pro Glu His Leu Ala
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Pro Gln Gln Asp Gln Val Pro Thr Gln Ala Gln Ser Gln Glu Gln Thr
 260 265 270

Ser Glu Lys Thr Gln Asp Gln Pro Gln Thr Trp Pro Gln Gly Ser Val
 275 280 285

Pro Pro Pro Glu Gln Ala Ser Gly Pro Ala Cys Ala Thr Glu Pro Gln
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Leu Ser Ser His Ala Ala Glu Ala Gly Ser Asp Pro Asp Lys Ala Leu
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 325 330 335

Ser Ala Gly Gly Leu Asp Leu Gly Glu Cys Glu Lys Arg Ala Gly Glu
340 345 350

Met Leu Gly Met Trp Gly Ala Gly Ser Ser Leu Lys Val Thr Ile Leu
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Gln Ser Ser Asn Ser Arg Ala Phe Asn Thr Thr Pro Leu Thr Ser Gly
370 375 380

Pro Arg Pro Gly Asp Ser Thr Ser Ala Thr Pro Ala Ile Ala Ser Thr
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Pro Ser Lys Gln Ser Leu Gln Phe Phe Cys Tyr Ile Cys Lys Ala Ser
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Ser Ser Ser Gln Gln Glu Phe Gln Asp His Met Ser Glu Ala Gln His
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Gln Gln Arg Leu Gly Glu Ile Gln His Ser Ser Gln Thr Cys Leu Leu
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